

D. Fitzgerald

1647

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/215,212

DATE: 08/08/2000
TIME: 14:51:41

Input Set : A:\TEPPER1A.txt
Output Set: N:\CRF3\08082000\I215212.raw

2 <110> APPLICANT: TEPPER, Mark
3 CUNNINGHAM, Mark
4 SHERRIS, David
5 EL TAYAR, Nabil
6 MCKENNA, Sean
W--> 7 <120> TITLE OF INVENTION: IFNAR2/IFN COMPLEX
W--> 8 <130> FILE REFERENCE: TEPPER1A.SEQ
W--> 9 <140> CURRENT APPLICATION NUMBER: 09/215,212
10 <141> CURRENT FILING DATE: 1998-12-18
11 <150> PRIOR APPLICATION NUMBER: 60/068,295
12 <151> PRIOR FILING DATE: 1997-12-19
W--> 13 <160> NUMBER OF SEQ ID: 15
14 <170> SOFTWARE: PatentIn Ver. 2.0

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

240 <210> SEQ ID NO: 15
241 <211> LENGTH: 8
242 <212> TYPE: PRT
243 <213> ORGANISM: Artificial Sequence
W--> 244 <220> FEATURE:
245 <223> OTHER INFORMATION: Description of Artificial Sequence: C terminal human
246 sIFNAR2 directly connected to N terminal human IFNbeta
248 <400> SEQUENCE: 15
249 Glu Ser Glu Phe Ser Met Ser Tyr
250 1 5
E--> 251 1

VERIFICATION SUMMARY DATE: 08/08/2000
PATENT APPLICATION: US/09/215,212 TIME: 14:51:42

Input Set : A:\TEPPER1A.txt
Output Set: N:\CRF3\08082000\I215212.raw

L:7 M:283 W: Missing Blank Line separator, <120> field identifier
L:8 M:283 W: Missing Blank Line separator, <130> field identifier
L:9 M:283 W: Missing Blank Line separator, <140> field identifier
L:13 M:283 W: Missing Blank Line separator, <160> field identifier
L:15 M:283 W: Missing Blank Line separator, <210> field identifier
L:19 M:283 W: Missing Blank Line separator, <220> field identifier
L:21 M:283 W: Missing Blank Line separator, <400> field identifier
L:24 M:283 W: Missing Blank Line separator, <210> field identifier
L:36 M:283 W: Missing Blank Line separator, <210> field identifier
L:40 M:283 W: Missing Blank Line separator, <220> field identifier
L:50 M:283 W: Missing Blank Line separator, <210> field identifier
L:54 M:283 W: Missing Blank Line separator, <220> field identifier
L:57 M:283 W: Missing Blank Line separator, <400> field identifier
L:62 M:283 W: Missing Blank Line separator, <210> field identifier
L:66 M:283 W: Missing Blank Line separator, <220> field identifier
L:69 M:283 W: Missing Blank Line separator, <400> field identifier
L:79 M:283 W: Missing Blank Line separator, <220> field identifier
L:82 M:283 W: Missing Blank Line separator, <400> field identifier
L:86 M:283 W: Missing Blank Line separator, <210> field identifier
L:90 M:283 W: Missing Blank Line separator, <220> field identifier
L:93 M:283 W: Missing Blank Line separator, <400> field identifier
L:101 M:283 W: Missing Blank Line separator, <220> field identifier
L:104 M:283 W: Missing Blank Line separator, <400> field identifier
L:111 M:283 W: Missing Blank Line separator, <210> field identifier
L:115 M:283 W: Missing Blank Line separator, <220> field identifier
L:118 M:283 W: Missing Blank Line separator, <400> field identifier
L:123 M:283 W: Missing Blank Line separator, <210> field identifier
L:127 M:283 W: Missing Blank Line separator, <220> field identifier
L:136 M:283 W: Missing Blank Line separator, <210> field identifier
L:140 M:283 W: Missing Blank Line separator, <220> field identifier
L:143 M:283 W: Missing Blank Line separator, <400> field identifier
L:152 M:283 W: Missing Blank Line separator, <220> field identifier
L:155 M:283 W: Missing Blank Line separator, <400> field identifier
L:165 M:283 W: Missing Blank Line separator, <220> field identifier
L:175 M:283 W: Missing Blank Line separator, <210> field identifier
L:179 M:283 W: Missing Blank Line separator, <220> field identifier
L:182 M:283 W: Missing Blank Line separator, <220> field identifier
L:244 M:283 W: Missing Blank Line separator, <220> field identifier
L:255 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:15